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REFERENCE AUTHORS TITLE JOURNAL	SOURCE	VERSION KEYWORDS	RESULT 1 AX395268 LOCUS DEFINITION ACCESSION		44.5	43	<b>4</b> 1						υ Ω Ω		29										14							×ω	21	No.	1
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Direct Submission
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Immunology, NIAID, NIH,
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Direct Submission
Submitted (16-NOV-1999)
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Submitted (13-JAN-1998)
Immunology, NIAID, NIH,
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Kim,M.G., Chen,C., Lyu,M.S.,
Schwartz,R.H.
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Mus musculus epithin mRNA,
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T.M.G., Chen,C., Cho,E., P.
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,M.G., Chen,C., Cho,E.,
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EWPMQVSLHALGQGHLCGASLISPDKLVSAARGFQDDKNFKSDYTMMTAFLGLLDQS
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KRSASGVQELKLKRIITHPSFNDFTFDYDIALLELEKSVEYSTVVRPICLPDATHVFF
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                                                                                                                                                                                                                                                                                                                                                                                                                               cell_type="stromal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          strain="C.B.17scid"
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CTGAGCTGGGTGAAGATGATGCTGTCCCGGAGAGCTGCTTCCCAACTGTCATTGAGCTCCC
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                                                                                                                                                                                                                                                                 TTGCGTGCTAGCGGCCCAGCCTGGGGGCAAGGGTTTGATGGCAGCCTTCCCCCCTCTAGCC
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old, gross tissue."
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/db\_xref="taxon:10090" /clone="MGC:7395 IMAGE:3488059" /tissue\_type="Mammary tumor, WA' old, gross tissue,"

WAP-TGF alpha

model.

J

/organism="Mus musculus" /mol\_type="mRNA" /strain="C57BL/6J"

note="Vector: pCMV-SPORT6"

/lab\_host="DH10B"

Location/Qualifiers

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Klausner, R.D., Collins, F.S., Wagner, L., Shenmen, C.M., Schuler, G.D., Klausner, R.D., Collins, F.S., Wagner, L., Shenmen, C.M., Schuler, G.D., Altschul, S.F., Zeeberg, B., Buetow, K.H., Schaefer, C.F., Bhat, N.K., Hopkins, R.F., Jordan, H., Moore, T., Max, S.I., Wang, J., Hsieh, F., Diatchenko, L., Marusina, K., Farmer, A.A., Rubin, G.M., Hong, L., Stapleton, M., Soares, M.B., Bonaldo, M.F., Casavant, T.L., Scheetz, T.E., Brownstein, M.J., Usdin, T.B., Toshiyuki, S. Carninci, P., Prange, C., Raha, S.S., Loquellano, N.A., Peters, G.J., Abramson, R.D., Mullahy, S.J., Bosak, S.A., McEwan, P.J., McKernan, K.J., Malek, J.A., Gunaratne, P.H., Richards, S., Worley, K.C., Hale, S., Garcia, A.M., Gay, L.J., Hulyk, S.W., Villalon, D.K., Muzny, D.M., Sodergren, E. J., Lu, X., Gibbs, R.A., Fahey, J., Helton, E., Ketteman, M., Madan, A., Rodrigues, S., Fahey, J., Helton, E., Ketteman, M., Madan, A., Rodrigues, S., Sanchez, A., Whiting, M., Madan, A., Young, A.C., Shevchenko, Y., Bouffard, G.G., Blakesley, R.W., Touchman, J.W., Green, E.D., Bouffard, Y.S., Krzywinski, M.I., Skalska, U., Smailus, D.E., Dickson, M.C., Rodriguez, A.C., Grimwood, J., Schmutz, J., Myers, R.M., Butterfield, Y.S., Krzywinski, M.I., Skalska, U., Smailus, D.E., Schnert, A., Schein, J.E., Jones, S.J. and Marra, M.A. Generation and initial analysis of more than 15,000 full-length human and mouse cDNA, sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Lothar Hennighausen Ph.D., Robin Humpl
CDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
Clone distribution: MGC clone distribution information can be for through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Series: IRAK Plate: 7 Row: i Column: 16 This clone was selected for full length sequencing because it passed the following selection criteria: Hexamer frequency ORF
                                                                                                                                                                                                                                                                                                                                     DNA Sequencing by: Sequencing Group at the Stanford Human Genome Center, Stanford University School of Medicine, Stanford, CA 943 Web site: http://www-shgc.stanford.edu Contact: (Dickson, Mark) mcd@paxil.stanford.edu Dickson, M., Schmutz, J., Grimwood, J., Rodriquez, A., and Myers,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Direct Submission
Submitted (27-MAR-2001) National Institutes of Health, Mammalian Submitted (27-MAR-2001) Cancer Genomics Office, National Cancer Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Strausberg, R
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AGKALWVTGMGHTKEGGTGALILOKGGEIRVINQTTCBDLMAPQQITFRMMCVGFLSGGV
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173 2740
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1640. .1747
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                                                                                                                                                                                                                                                                                                                                                                                                       /note="LDLa; Region: Low Density Lipoprotein Receptor Class A domain, a cysteine-rich repeat that plays a central role in mammalian cholesterol metabolism"
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domain class A"
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domain class A"
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Pred. No. 0;
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                         actateregaearcaacegegaeaactactecegreacaegercccaetrieregreaca
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2340 2450	281 CTGCTGGCAAGGCCATCTGGGTCACAGGCTGGGGGCACACAAAAGAGGGAGG
2280 2390	221 CGGTGGAGTACAGCACCGTCGTGGCGCCCCATCTGCCTGC
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2160 2270	2101 ACCAGAGCAAGCGCAGTGCCTCTGGGGTGCAGGAGCTGAAGCTCAAACGTATCATCACCC
2100	2041 ATGACAAAATTTCAAGTACTCAGACTACACGATGTGGACGGCCTTCCTGGGTCTGCTGG
2040	1981 TGTGTGGGGCCTCGCTCATCTCCTGACTGGCTGGTTGTTGTTGTTTTCAGG
1980 2090	1921 ATGCGGACGAGGGCGAGTGGCCCTGGCAGGTGAGCCTCCACGCCCTGGGCCAGGGCCACT
1920 2030	1861 AAAACTGTGACTGTGGGCTGCGATCCTTTACCAAACAGGCTCGCGTGGTTGGT
1860	1801 TGAGCAAGGGCAACCCTGAGTGTGATGGGAAGACGGACTGTAGCGATGGCTCCGATGAGA
1800	0-0
1740 1850	1681 AGAGCCAGAAGTGTAATGGGAAGGACAACTGTGGAGATGGGTCTGACGAGGCTTCATGTG
1680 1790	1621 AGGAGGCTGCAGCTGTCCTGCTGGGAGTTTCAAGTGTTCCAATGGGAAGTGTCTCCCTC
1620 1730	1561 TCTGCAAGCCCCTCTTCTGGGTCTGTGACAGTGTCAACGACTGTGGGGACGGAAGTGACG
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1500	1441 AGACTGGACGGTGCATCCGAAAGGAACTGCGCTGCGACGGCTGGGCAGACTGCCCGGATT
1440	CTGAGTACCT           CTGAGTACCT
1380 1490	.321 GCAACAGCAGCAAGATTACAG 

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ACCESSION AB04911
VERSION AB04911
KEYWORDS RATTUS
SOURCE BUARTY
ORGANISM RATTUS
EUARTY
ORGANISM RATTUS
AUTHORS INOUE, I
AUTHORS INOUE, I
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TITLE DIRECT
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RESULT 4
AB049189
AB049189
AB049189
AB049189
ABOTOUS
DEFINITION Rattus norvegicus mRNA for membrane bound arginine specific serine
protease, complete cds.
ACCESSION AB049189
VERSION AB049189.1 GI:10336526
KEYMORDS
SOURCE
ORGANISM Rattus norvegicus (Norway rat)
CORGANISM Rattus norvegicus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammala; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;

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I (sites)

Inoue,H., Takahashi,K. and Kishi,K.

membrane-bound arginine specific serine protease

Published Only in DataBase (2000)

(bases 1 to 3174)

Inoue,H., Takahashi,K. and Kishi,K.

Direct Submission

Submitted (22-SEP-2000) Hideshi Inoue, Tokyo University of Pharmacy Submitted (22-SEP-2000) Hideshi Inoue, Tokyo 11492-1 Horinouchi, Hachiojishi, Tokyo 192-0392, Japan (B-mail:hinoue@ls.toyaku.ac.jp, Tel:81-426-76-7153, Fax:81-426-76-7157)

Location/Qualifiers

FEATURES

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     TCAAGTACAACTCCCGGCTAGAGAACATGAATGGCTTTTGAGGAGGTGTGGAGTTCCTGC
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TITLE JOURNAL	2461 CCTGCCAGGGTGACTCTGGTGGCCCCTTGTCAAGCGCGGAGAAAGATGGGCGAATGTTCC 2520	γQ
MEDLINE PUBMED REFERENCE AUTHORS	2401 TGCCGCAGCAGATCACCCCACGAATGATGTGTGTGGGTTTCCTCAGTGGGGGTGTGGACT 2460	dg VQ
TITLE	2341 CGCTGATCCTGCAGAAGGGTGAGATCCGTGTCATCAACCAGACCACCTGTGAGGACCTCA 2400	Db
REFERENCE AUTHORS	w N	D Qy
SOURCE ORGANISM	221 C	90 VQ
ACCESSION VERSION	2161 ACCCTTCCATGATTCACCTTCGACTATGACATCGCCTTGCTGGAGCTGGAGAAGT 2220	B 8
RESULT 5 AB037898 LOCUS	2101 ACCAGAGCAAGCGCAGTGCCTCTGGGGTGCAGGAGCTGAAGCTCAAACGTATCATCACCC 2160	P &
Db 31	S ATGAGACAATTTTCAAGTACTCAGACCACCATGTGGACAGCCTTCCTCGGTCTGCTGG	Db
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	1861 AAAACTGTGAGTGTGGGCTGCGATCCTTTACCAAACAGGCTCGCGTGGTTGGT	Ş
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Дb 2	1625 TCTGCAAGCCCTCTTCTGGGTCTGTGACAGTGTCAACGACTGTGGGGATGGCGATG 1684	Db
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Rattus norveg complete cds. AB037898 AB037898.1 GI:9650963 membrane bound serine protease; Rattus norvegicus (Norway rat) Rattus norvegicus norvegicus MBSP mRNA for membrane bound serine protease, mRNA linear ROD 20-OCT-2001

MT-SP1

Eukaryota; Metazoa; Mammalia; Eutheria; Rattus Chordata; Rodentia; Tsuzuki,S., Craniata; Vertebrata; I Sciurognathi; Muridae; Hitomi, Y., Iwanaga, T. Euteleostomi;
; Murinae;

Satomi,S., Yamasaki,Y., Ts Fushiki,T. A role for membrane-type s epithelial turnover Biochem. Biophys. Res. Com 21458307 11573963 2 (bases 1 to 2568) Commun. 287 serine protease (MT-SP1) (4), 995-1002 (2001) in intestinal and

(bases 1 to 2568)

Tsuzuki,S.

Tsuzuki,S.

Direct Submission

Direct Submission

Submitted (26-JAN-2000) Satoshi Tsuzuki, Kyoto University, Graduato Submitted (26-JAN-2000) Satoshi Tsuzuki, Kyoto University, Graduato Submol of Agriculture, Division of Food Science and Biotechnology, Graduate

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/db_xref="GI:9650964"
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RS Strausberg R.L. Feingold, E.A., Grouse, L.H., Derge, J.G.,
Klausner, R.D., Collins, F.S., Wagner, L., Shenmen, C.M., Schuler, G.D.,
Altschul, S.F., Zeeberg, B., Buetow, K.H., Schaefer, C.F., Bhat, N.K.,
Hopkins, R.F., Jordan, H., Moore, T., Max, S.I., Wang, J., Hsieh, F.,
Diatchenko, L., Marusina, K., Farmer, A.A., Rubin, G.M., Hong, L.,
Stapleton, M., Soares, M.B., Bonaldo, M.F., Casavant, T.L.,
Scheetz, T.E., Brownstein, M.J., Usdin, T.B., Toshiyuki, S.,
Carninci, P., Prange, C., Raha, S.S., Loquellano, N.A., Peters, G.J.,
Abramson, R.D., Mullahy, S.J., Bosak, S.A., McEwan, P.J.,
McKernan, K.J., Malek, J.A., Gunaratne, P.H., Richards, S.,
Worley, K.C., Hale, S., Garcia, A.M., Gay, L.J., Hulyk, S.W.,
Villalon, D.K., Muzny, D.M., Sodergren, E.J., Lu, X., Gibbs, R.A.,
Fahey, J., Helton, E., Ketteman, M., Madan, A., Rodrigues, S.,
Bouffard, G.G., Blakesley, R.W., Touchman, J.W., Green, E.D.,
Dickson, M.C., Rodriguez, A.C., Grimwood, J., Schmutz, J., Myers, R.M.,
Butterfield, Y.S., Krzywinski, M.I., Skalska, U., Smailus, D.E.,
Generation and inirial snalveis of force than 15 000 finil-locath
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Contact: nisc_mgc@nhgri.nih.gov
Akhter,N., Ayele,K., Beckstrom-Sternberg,S.M., Benjamin,B.,
Blakesley,R.W., Bouffard,G.G., Breen,K., Brinkley,C., Brooks,S.,
Blakesley,R.W., Bouffard,G.G., Breen,X., Gupta,J., Haghighi,P.,
Hansen,N., Ho,S.-L., Karlins,E., Kwong,P., Laric,P., Legaspi,R.,
Maduro,Q.L., Masiello,C., Maskeri,B., Mastrian,S.D., McCloskey,J.C.,
McDowell,J., Pearson,R., Stantripop,S., Thomas,P.J., Touchman,J.W.,
Tsurgeon,C., Vogt,J.L., Walker,M.A., Wetherby,K.D., Wiggins,L.,
Young,A., Zhang,L.-H. and Green,E.D.
                                                                                                                                                                                                                                                                                                                                              Clone distribution: MGC clone distribution information can be for through the I.M.A.G.E. Consortium/LLML at: http://image.llnl.gov Series: IRAK Plate: 64 Row: k Column: 15
This clone was selected for full length sequencing because it passed the following selection criteria: Similarity but not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                   passed the following identity to protein.
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Submitted (07-MAY-2002) National Institutes of Health, Mammalian Sene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaithersburg, Maryland;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA Sequencing by: National Institutes of Health Intramural Sequencing Center (NISC),
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/organism="Homo sapiens"
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/db_xref="raxon:9606"
/clone="MGC:40392 IMAGE:5213189"
/tissue_type="Blood, adult leukocytes"
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              TCAAGTACAACTCCCGGCACGAGAAAGTGAATGGCTTGGAGGAAGGCGTGGAGTTCCTGC
                                               TCRAGTACACTCCCGGCTAGAGAACATGAATGGCTTTGAGGAGGTGTGGAGTTCCTGC
                                                                                                                                                                CCATGGGTAGCAATCGGGGGCCGCAAGGCCGGAGGGGGCTCTCAGGACTTCGGCGCGGGAC
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DPCPGQFTCRTGRCIRKELRCDGWADCTDHSDELNCSCDAGHQPTCKNKFCKPLFWVC
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DSCQGDSGGPLSSVEADGRIFQAGVVSWGDGCAQRNKPGVYTRLPLFRDWIKENTGV"
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PRMSSCGGRLRKAQGTFNSPYYPGHYPPNIDCTWNIEVPNNGYKVRFKFFYLLEPGV
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/lab_host="DH10B"
/note="Vector: pCMV-SPORT6"
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/note="synonyms: SNC19,
HAI, MT-SP1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1690. .1800
/note="LDLa; Region: Low-density
domain class A"
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/db_xref="CDD:smart^^^^^
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/note="CUB; Region: CUB domain"
/db_xref="CDD:pfam00431"
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C12N1/37,G01N33/53,G01N33/577/C12P21/08,A61K49/02,C12N15/00,
PC C12N5/00
CC Matriptase, serine protease and application of ""
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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 3149)

1 (ckson,R.B., Lin,C., Johnson,M., Wang,S. and Enye,I.
Matriptase, serine protease and application of the same
GEORGETOWN UNIVERSITY

OS Homo sapiens (human)

JP 2002539033-A/2

PD 19-NOV-2002

PF 10-MAR-2000 JP 2000603721

PF 10-MAR-2000 JP 2000603721

PF 10-MAR-1999 US 60/124006

PI ROBERT B DICKSON,CHEN-YONG LIN,MICHAEL JOHNSON,SHAOMENG WANG,
PI ISTVAN ENYEDY

PC A61843/00,A61835/78,A61851/00,A61P1/00,A61P17/00,A61P35/00, PC
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                                                                      3 (bases 1 to 3149)
Oberst, M.D., Dickson, R.B. and Lin, C.Y.
Direct Submission
Submitted (30-DEC-1999) Lombardi Cancer Center,
University Medical Center, 3970 Reservoir Road
         Sequence update by submitter On Dec 30, 1999 this sequence
                                                                                                                                                                                                                    Submitted (06-JAN-1999) Lombardi
University Medical Center, 3970 I
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J. Biol. Chem. 274 (26), 18231-18236 (1999)
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Direct Submission
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                             GCGAGTTGTAACATTGCCACCCCGAGCACGGGCACTGAAAATCCTTCGTGCTAACATCTGT
GCGCGTAGTCATGCTGCCCCCGCGGGGCGCGCTCCCTGAAGTCCTTTGTGGTCACCTCAGT
                                                                                     GTCTGAGTTCAGCATCCCGCAGCACCTGGTGGAGGAGGCCGAGCGCGTCATGGCCGAGGA
                                                                                                                       GTCAGAGTTCAGCATCCCCCCCACACCTGGCAGAAGAGGTTGATCGCGCCCATGGCTGTGGA
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KKVEKIGPGRWTVLAAVLIGLLLVILGIGFLYWHLQYNDVRVQKVFNGYMRITNENFV
DAYENSNSTEFVSLASKVKDALKLLYSGVPFLGFYHKEAVTAFSEGSVLAYYMSEFS
I POHLVEEAERVWALERVMLDPRAFSLKSFVTSVTAFFDBSKTYQRTODNSCSFGL
HARGVELMRFTTPGFFDDSPYPAHARCQWALRGDADSVLSLTFRSFDLASCDERGSDLV
TVYNTLSPMEPHALVQLCGTYPFSYNLTFHSSGVVTLITITNTWERRHFGFEATFFQL
FRMSSCGGRLKKAGGTFNSFYPGHYPEN IDCTWNIEVPNNQHVKVRFKFFYLLEPGY
PAGTCFKDYVEINGEKYCGERSOFVVTSNSNKITVRFHSDQSYTDTGFLAEYLSYDS
DPCFGQFTCRTGRCIRKELRCDGWADCTDHSDELNCSCDAGHQFTCKNKFCKPLFWVC
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DSVNDCGDNSDBGGSCFAQTFRCSNGKTCKSKQCCNAKDDCGDGSDEASCFKNVVT
CTKHTYRCLNGLCLSKONPECDGKEDCSDGSDEKDCRCRSPETSDDTOWTAFLGLHDQS
QRSAFGYQERLKKRISHFFFNDFTFDYDIALLELEKPAEYSSMVFTCLDBASHYFP
AGKAIWTGWGHTQYGGTGALILOKGEIRVINQTTCENLLEQQITPRMMCVGFLSGGV
DSCQGDSGGPLSSVEADGRIFQAGVVSWGDGCAQRNKEGCYTRLPLFRDMKKENTGV"
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/protein_id="AAD42765.2"
/db_xref="GI:6647302"
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/note="serine protease with trypsin-like
contains Clr/s and LDL receptor domains"
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/db_xref="taxon:9606"
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650 GRITGGCCTTCCCCATTGACCCCAAAACAGTCCCAAGGACTCCAGGACAACAGCTCCAGTTT 709 621 GRITGGCTTTCCCCACGTCCCAAAACAGTCCCCAAAACAGTCCAGGACAACAGCTCCAGTTT 692 710 TCCCTACCCTCCCAAGCCTTGCCCACGTCCCAAGCAGTCCCCAAGCAGTCCCCCAAGCTCCCCAAGCTCCACTCCCTCC
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Ωу 21	Qy 154 Db 128	Db e	Qy Db	Query Match Best Local Matches 216	ORIGIN					CDS		FEATURES Source	REFERENCE AUTHORS TITLE JOURNAL	JOURNAL MEDLINE	REFERENCE AUTHORS TITLE	SOURCE ORGANISM	DEFINITION ACCESSION VERSION KEYWORDS	RESULT 9 AF133086
	54 GCTTTGAGGAGGGTGTGGAGTTCCTGCCGAGCAATGCCAAGAAAGTGGAGAAAGCGAG 213	94 GGGGCTCTCAGGACTTCGGCGCGGGACTCAAGTACAACTCCCGGCTAGAGAACATGAATG 153 	₩ <b>~</b> ₩	ttch 60.7%; Score 1884.8; DB 9; Length 3120; al Similarity 82.8%; Pred. No. 0; 2164; Conservative 0; Mismatches 447; Indels 1; Gaps 1;	QKSAFGVQEKKIKKI ISHPFENDFTFDYD JALLEKPARYSSMYRPICLPDASHYFP AGKAIWVTGWGHTQYGGTGALILQKGEIRVINQTTCENLLPQQITPRMMCVGFLSGGV DSCQGDSGGPLSSVEADGRIFQAGVVSWGDGCAQRNKPGVYTRLPLFRDWIKENTGV"	DECEGRETICATION OF THE SELECTION OF THE	HARGYELMRETTPOEPDSPYPAHARCQWALRGDADSVLSLTFRSFDLASCDERGSDLV TVNITLS PMEPHALVQLCGTYPSSYNLTFHSQNVLLITLLINTERRHPGFBATFFQL PRMSSCGGRLRKAQGTFNSBYYPGHYPBNIDGTMNIEVPNNOQHVKVFRKFFYLLEPGV PAGTCPKDYVEINGEKYCGERSQFVVTSNSNKITVRFHSDGYTDTGFLABYLSYDSS	/ translation="WGSDRARKGGGGPKDPGGLKYNGSRHEKVNGLEEGVEFLPVNNVKYEKVEKHGPGRWVLAAVLIGLLIVLLGIGFLVMHLQYRDVRVQKVFNGYMRITNBRFVDAYBKSRGTEFVSLAKVKRALKLLYSGVPFLGPYHKESAVTAFSEGSVIAYYMSEFGLFVGLAKVKRALKLLYSGVPFLGPYHKESAVTAFSEGSVIAYYMSEFGLFVGLAKVKRALKLLYSGVPFUGYHKESAVTAFSEGSVIAYYMSEFGLFVDAKVFVDAFFTDSKTYQRTQNNSCSFGL	/codon_start=1 /product="membrane-type_serine_protease_1" /protein_id="AAR00109.1" /db_xref="GI:6002714"			California, San Francisco, 513 Parnas Francisco, CA 94143-0446, USA Location/Qualifiers	Craik, C.S.  Pharm Chem University	serine protease in epithelial cancer and normal trissue Proc. Natl. Acad. Sci. U.S.A. 96 (20), 11054-11061 (1999) 99432178	1 (bases 1 to 3120)  Takeuchi, T., Shuman, M.A. and Craik, C.S.  Reverse biochemistry: use of macromolecular protease inhibitors to dissect complex biological processes and identifications.	Homo sapiens (human) Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates: Catarrhini. Hominidae. Homo	Homo sapiens membrane-type serine protease 1 mRNA, complete cds. AF133086 AF133086.1 GI:6002713	
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1414 ACCCGTGCCCAGGATGTTCATGTGCAAGACTGGACGGTGCATCCGAAAGGAACTGCGCT 1473 1414	474 GCGACGGCTGGGCAGACTGCCCGGATTATAGTGATGAGGGTTACT	տ տ	594 T 568 T	654 AGTGTTCCAATGGGAAGTGTCTCCCTCA	714 GAGATGGGTCTGACGAGGCTTCATGTGACAGCGTGAATGTCGTCTCTTGCACCAAATATA 17 	774 CCTACCGCTGCCAAAATGGCCTCTGTCTGAGCAAGGGCAACCCTGAGTGTGATGGGAAGA 18	<b>ω</b> ω	954 GCCTCCACGCCCTGGGCCAGGCCAGTTGTG: 	014 TGGTCTCTGCAGCTCATTGCTTTCAGGATGA	0 0	134 AGCTGAAGCTC           108 AGCGCAGGCTC	194 ACATCGCCTTC            168 ACATCGCGCTC	2254 GCCTGCCTGATGCTACCCATGTCTTCCCTGCTGGCAAGGCCATCTGGGTCACAGGCTGGG 2313	2314 GGCACAAAAGAGGGAGGTACCGGAGCGCTGATCCTGCAGAAGGGTGAGATCCGTGTCA 237	2374 TCAACCAGACCACCTGTGAGGACCTCATGCCGCAGCAGATCACCCCACGAATGATGTG 2433
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AB030036.1
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Molecular cloning of a novel transmembrane serine protease expressed in human prostate
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Yamaguchi, N. and Mitsui, S.
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/protein_id="BAB20376.1"
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S	Qy 2750 GACTGTGAACTGCATCCTTAGGACTCAGAGT 2780	0-0 0-0 4-4		QY 2270 CCATESTETTCCCTGCTGGCAAGGCCATCTGGGTCACAGGCTGGGGCACACAAAAGAGG 2329
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	2345 ACCTGCGAGAACCTCCTGCCGCAGCAGATCACGCCGCGCATGATGTGCGCTGGGCTTCCTC  2445 AGTGGGGGTGTGGACTCCTGCCAGGGTGACTCTGGTGGCCCCTTGTCAAGCGCGGAGAAA  2445 AGTGGGGGTGTGGACTCCTGCCAGGGTGACTCTGGTGGCCCCTTGTCAAGCGCGGAGAAA  2405 AGCGGCGGGTGGACTCCTGCCAGGGTGATTCCGGGGGAACCCCTTGTCCAGCGTGAAGGAAC  2405 AGCGGCGGGTGGACTCCTGCCAGGGTGATCCCGGGTGAAGGCTCAGAGGAAC  2505 GATGGGCGAATGTTCCAGGCTGGTGTGGTGAGCTGAGGCTGAAGGAAC  2465 GATGGGCGAATCTTCCAGGCCGGTGTAGTTCGGGAACCTGGATCAAAGAGCACACTGGG  2565 AAGCCAGGCGTGTACACAAAGGCTCCCTGTAGTTCGGGACTGGATCAAAGAGCACACTGGG  2565 AAGCCAGGCGTGTACACAAAGGCTCCCTGTAGTTCGGGACTGGATCAAAAGAGCACACTGGG  2565 AAGCCAGGCGTGTACACAAAGGCTCCCTGTAGTTCCGGGACTGGATCAAAGAGCACACTGGG		985 985 085 045 145	1625 1685 1685 1785 1785 1785 1985 1886 1886 1886 1886 1886 1886 1886 18

QY 705 AGTITITGCCCTGCATGGTGCAGGTGCAGCAGCGTTCACCTCTGGCTTCCCC 764	2 7	GCTAACA 6	N O	QY 465 GGTCCCTACCACAAGAAGTCGGCTGTAACTGCCTTCAGTGAGGGCAGTGTCATCGCCTAC 524	OY 405 AGCCTGGCCAGCCAGGTGAAGGAGGCGCTGAAGCTGCTGTACAATGAAGTCCCTGTCCTG 464	QY 345 AGGATCACAAATGAGATCTTTCTGGATGCGTATGAGAACTCCACCTCCACAGAGTTTATC 404	QY 285 CTGGTGTGGCACTTCCATTATCGGAATGTGCGGGTTCAAAAAGTCTTCAATGGCCATCTG 344	Qy 225 TGGGTGGTGGCAGTGCTGTTCAGCTTCCTCTTGCTCTCATGGCTGGC	, N N	Qy 105 GACTTCGGCGGGGACTCAAGTACAACTCCCGGCTAGAGAACATGAATGGCTTTGAGGAG 164	OY  45 GATICGGACCGCCAAAACCATGGGTAGCAATCGGGCCCGCAAGGGCGGAGGGGGCCCGAAG 3084  Db 3143 GAGCGGCCTCGGGGTACCATGGGGAGCGATCGGGCCCGCAAGGGCGGAGGGGGCCCGAAG 3084	60.6%; Similarity 81.2%; 3; Conservative	rce 13147 /organism="unk: /mol_type="una	TITLE TADG-15: an extracellular serine protease overexpressed in carcinomas  JOURNAL Patent: US 6649741-A 18 18-NOV-2003;  FRATURES Location/Qualifiers	Unknown.  SM Unknown.  Unclassified.  1 (bases 1 to 3147)  CO'Brien T.J. and Tanimoto.	NITION Sequence 18 from patent SSION AR430938 I GI:40192680 ORDS	3147 bp	
Db 1403 CTCAATGGGCTCTGACCAAGCGCAAGCCCTGAGTGTGACGGGAAGGACTGTAGC 1344	Db 1463 GACGAGGCCTCCTCCCCAAGGTGAACGTCGTCAAACACACCTACCGCTGC 1404  OY 1785 CAAAATGGCCTCTGTCTGAGGAAAGGCAACCCTGAGGGAAGACGAAGACGAAGACGAACTGTAGC 1844	Db 1523 GGGAAGTGCCTCTGAAAAGCCAGCAGTGGAAAGGACGACTGTGGGGAACGGGTCC 1464  QY 1725 GACGAGGCTTCATGTGAACAGCGTGAATGTCGTCTTGCACAAATATACCTACC						Db 1883 CAGTTCGTCGTCACCAGCAACAAGATCACAGTTCGCTTCCACTCAGATCAGTCC 1824  Oy 1365 TACACGGACACGGGTTCCTAGCTGAGTACCTCTCCTACGACTCCAACGACTCGTGCCA 1424	Db 1943 GCCACCTGCCCCAAGGACTACGTGGAGATCAATACTGCGGAAGAGAGGTCC 1884  Oy 1305 CAGTTTGTGGTGAGCAGCAACAGCAGCAAGATTACAGTCCACTTCTGATCACTCG 1364	Db 2003 AACCAGCATGTGAAGGTTCAAATTCTTCTACTGTGGAGCCCGGCGTGCCTGCG 1944  Qy 1245 GGCTCCTGCACCAAGGACTATGTGGAGACAGAGAGAAGAAGAGGTCC 1304	Db 2063 TACTACCCAGGCCACTACCCAACATTGACTGCACATGGAACATTGAGGTGCCCAAC 2004  Qy 1185 AACCGGAACGTGAAGGTGCACTTCAAACTCTTCTATCTGGTGGACCCCAACGTACCAGTG 1244	2123 CCTAGGATGAGCAGCTGTGGAGGCCGCTTACGTAAAGCCCAGGGGACATTCAACAGCCCC  1125 TACTATCCAGGCCACTACCGCCCAACATCAACTGCACATGGAATATCAAGGTGCCCAAC	Oy 1005 ACGCIGATIACCARLACTGACCAGCGATCCCGGCTTTGAGGCCACCTTCTCCAGCTG 2124  Db 2183 ACACTGATAACCAACACTGAGGCGGCTTTTGAGTGACCCCAAGGGACATTTAGCAGCCC 1124  Oy 1065 CCCAAGATGAGCAGCTGTGGCGGCTTTTTGAGTGACCCCAAGGGACATTTAGCAGCCC 1124	945 2243	885 2303	QY 825 GTGCTGAGCCTCACCTTCGAAGCTTTGATGTCGCTCCTGTGATGAGCATGACCATGGCAGGAC 884	Oy 765 AACAGTCCCTACCCGGCGCATGCCCAGTGGGTCCTGCGGGGGGACGCCGACTCT 824	Db 2483 AGCTTTGGCCTGCACGCCCGGGTGTGGAGCTGATGCGCCTTCACCACGCCCGGCTTCCCT 2424

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